AMENDMENTS

Listing of Claims

The following listing of claims replaces all previous listings or versions thereof:

- 1. (Currently amended) A method for detecting endotoxin, comprising the steps:
 - a) incubating a sample with [[a]]an isolated bacteriophage tail protein, and
 - b) detecting endotoxin bonded to bacteriophage tail proteins.
- (Currently amended) The method according to claim 1, further comprising after step a) and prior to step b) the additional step of:
 - a') separating the-bacteriophage tail protein-endotoxin complexes from the sample.
- (Previously presented) The method according to claim 1, wherein detection comprises spectroscopic methods.
- (Currently amended) A method for removing endotoxin from a sample, comprising the steps:
 - incubating a sample with or bringing a sample in contact with <u>isolated</u> bacteriophage tail proteins which are immobilised on a permanent carrier, nonspecifically or directed,
 - separating the-bacteriophage tail protein-endotoxin complex from the sample.
- (Previously presented) The method according to claim 4, wherein steps a) and b) are implemented in a chromatography column throughflow method.
- 6. (Previously presented) The method according to claim 4, wherein the permanent carrier comprises filtration media, glass particles, magnetic particles, centrifugation materials, sedimentation materials or filling materials for chromatography columns.

- (Previously presented) The method according to claim 4, the bacteriophage tail proteins being immobilised on the permanent carrier via coupling groups.
- (Previously presented) The method according to claim 7, the coupling group being a lectin, receptor or anticalin.
- (Currently amended) The method according to claim 7, wherein the coupling group comprises streptavidin or avidin and the bacteriophage tail proteins [[is]]<u>are</u> coupled with biotin or a Strep-tag.
- (Currently amended) The method according to claim 4, the bacteriophage tail proteins being are immobilised on the permanent carrier covalently via chemical bonds.
- (Previously presented) The method according to claim 1, wherein the bacteriophage tail
 protein comprises a Strep-tag or a His-tag.
- (Previously presented) The method according to claim 11, wherein the tag comprises an amino acid sequence according to SEQ ID NO. 5, 6 or 7.
- 13. (Currently amended) The method according claim 11, wherein the p12 protein of the phage T4 is used as the bacteriophage tail protein.
- 14. (Previously presented) The method according to claim 1, wherein the Ca^{2+} concentration of the incubation comprises 0.1 μ M to 10 mM and the Mg^{2+} concentration comprises 0.1 μ M to 10 mM.
- 15. (Currently amended) The method according to one-of-the-claim 1, marked endotoxin being displaced from the bond with [[a]]the bacteriophage tail protein and the marked endotoxin being subsequently detected.
- 16. (Previously presented) The method according to claim 1, wherein the bacteriophage tail protein comprises a Strep-tag or a His-tag.
- 17. (Previously presented) The method according to claim 16, wherein the tag comprises an amino acid sequence according to SEQ ID NO. 5, 6 or 7.

18. (Currently amended) The method according claim 16, wherein the p12 protein of the phage T4 being used as the bacteriophage tail protein.